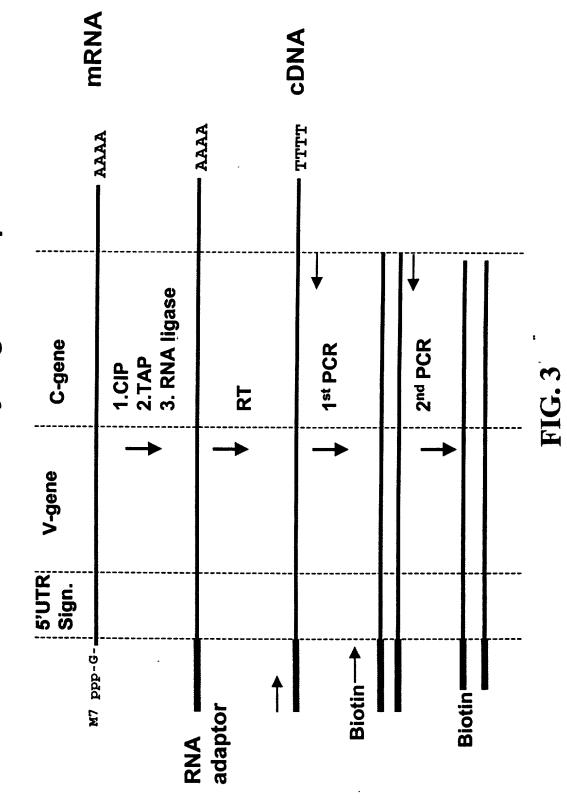


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mRNA 1ST STRAND DNA mRNA 1ST STRAND DNA dsDNA IMMOBILIZABLE ds cDNĄ C-SPECIFIC PRIMER www.www.www.www.www.www Y RE SITE CAPTURE ON STREPTAVIDIN BEADS minime. CCC production for the production of the UTR::SIGNAL VARIABLE REVERSE TRANSCRIPTION REVERSE TRANSCRIPTION *mmmmmm* AMPLIFY VL GENES WITHOUT USING VL SEQUENCES SMART PCR 4 ٥ $\mathbf{\omega}$ ပ ш

RACE non-biased antibody V-gene amplification



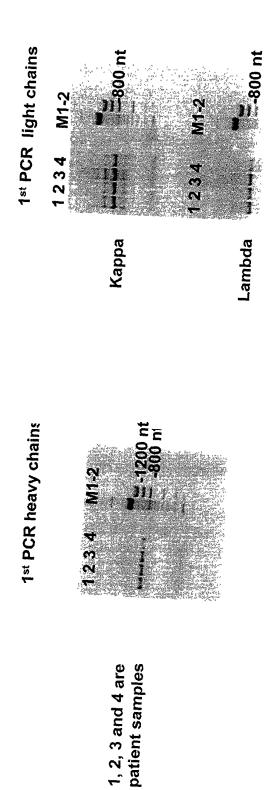
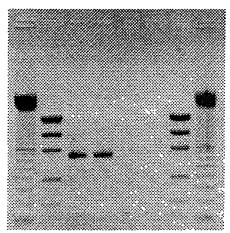


FIG. 4

4 5 6 1



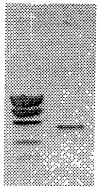
Gel analysis of PCR product from extender-kappa amplification
Approx. 75ng/5μl → 15ng/μl

- 1 100bp 2 LDM

- 3 50ng template
 4 10ng template
 5 ssDNA unligated
 6 negative control
- 7 LDM
- 8 100bp

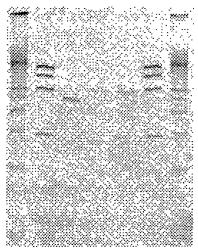
FIG. 5

1 2



Gel purified PCR product from extender-kappa amplification Concentration : $\pm 35 ng/\mu l$

- 1 LDM
- 2 1μl purif.

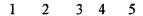


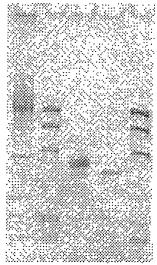
Gel-analysis of digested κ-ssDNA

- 1μl digested ssDNA ≈ 8ng ssDNA

 Total volume of 50μl = 400ng ssDNA

 400ng ssDNA available for ligation of the bridge-extenders
- 1 100bp
- 2 LDM
- 3 1μl ssDNA pure
 4 4μl beads after dig.
 5 8μl beads after dig.
- 6 LDM
- 7 100bp





Gel analysis of extender – cleaved kappa ligation 20ng/5µl eluted material → 4ng/µl

- 1- 100bp
- 2 LDM
- 3 Ligationmix, 4µl 4 Unligated ssDNA
- 5 LDM

FIG. 8

Cleavage and ligation Kappa light chains

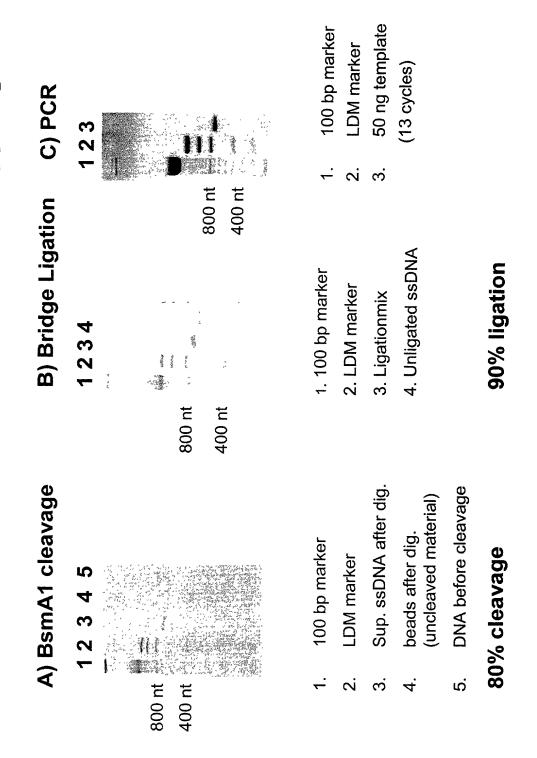


FIG. 9

VH-CDR1

1 Y 1 M 1

VH-CDR2

2 I 2 3 S G G 1 T 1 YADSVKG

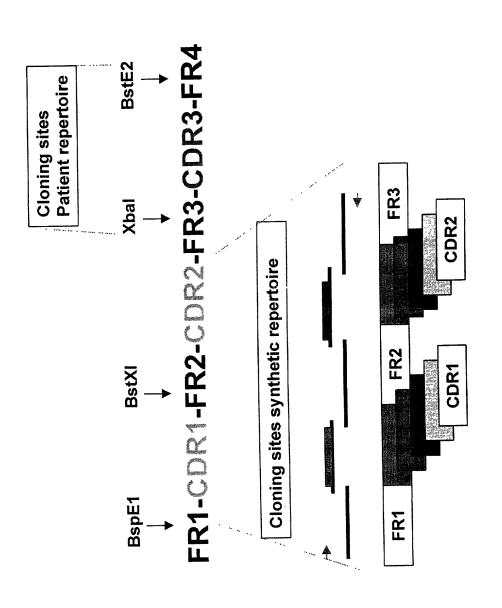


FIG. 11

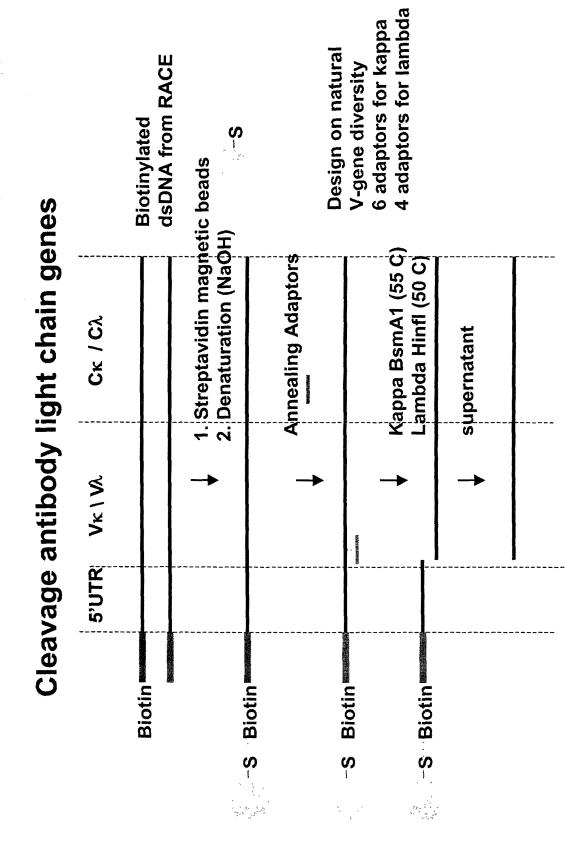


FIG. 12A

Ligation of cleaved light chains

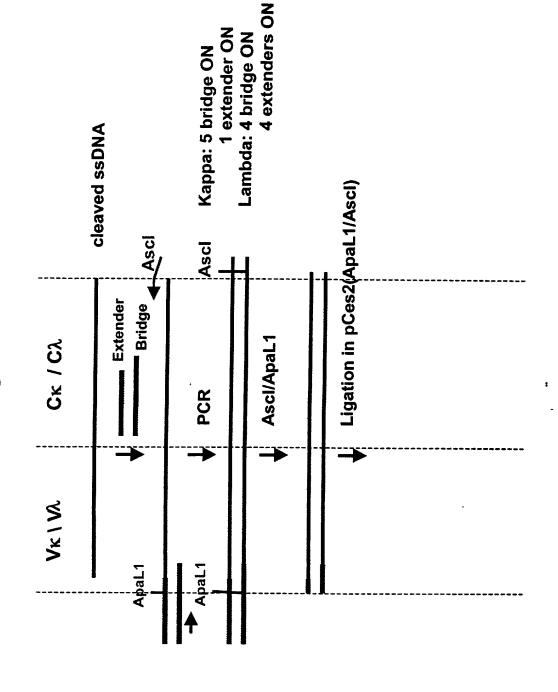


FIG. 12B

Figure 3: Cleavage and ligation lambda light chains

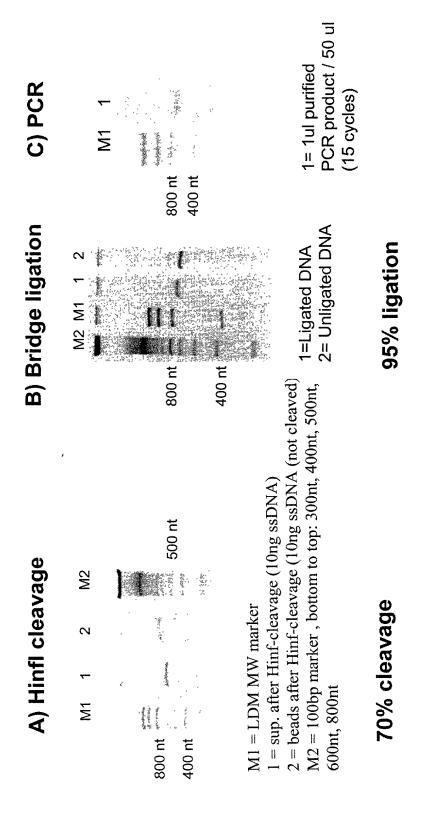


FIG. 13

CJ cleavage heavy chain

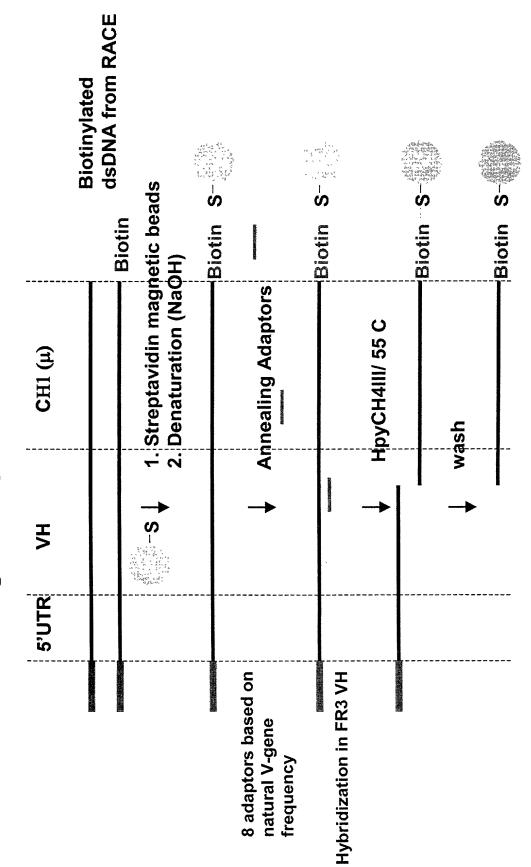


FIG.14A

Ligation heavy chain CDR3 diversity

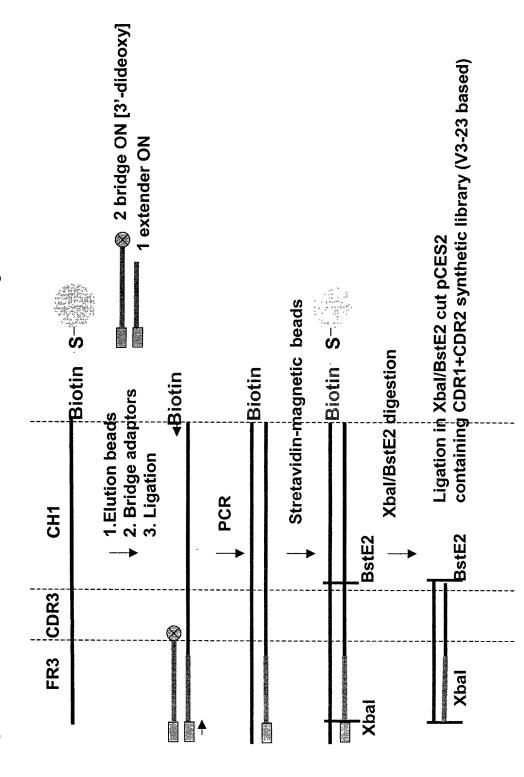
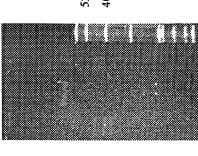


FIG. 14B

Cleavage and ligation Heavy Chain

B) PCR

A) HpyCH4III cleavage



527 nt

400 mt

500 bp

- 2 = 5ul/100ul PCR product 20 cycles; sample A

1 = NEB 100bp ladder

1 = Cleaved DNA eluted from PN column 2 = Beads after Hpy CH4III digestion

3 = Supernatant after cleavage 4 = MspI digest of pBR322

- 3 = 5ul/100ul PCR product 20 cycles; sample B
- 4 = no template

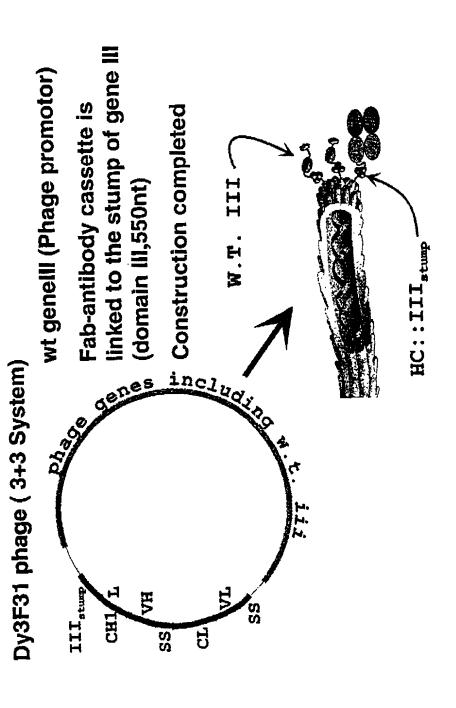


FIG. 16

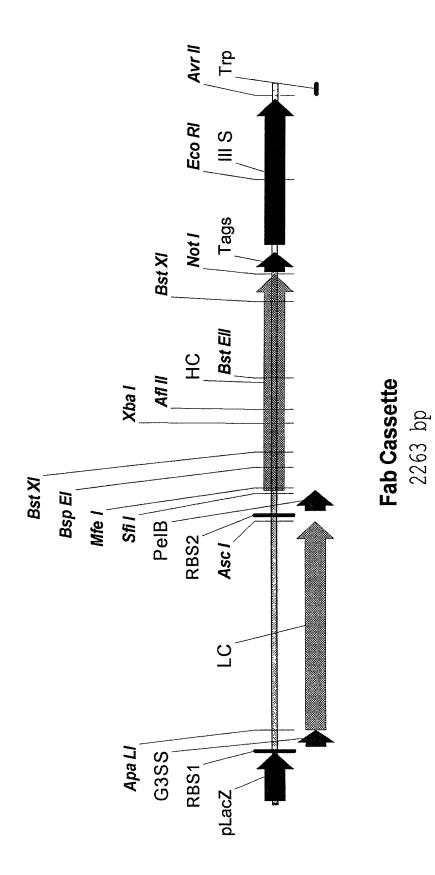


FIG. 17

1. Annealing

3. PCR

2. Ligation

PCRpr.: 5'CCTCGACAGCGAAGTGCA CAG-3'

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5, -

-ApaLI-

Q S A

AA-VL

S

1 +5

3. PCR

PCRpr.: 5'-CCTCTGTCACA GTGCA CAA GAC-3'

1. Annealing 5'-XXX-XXX X-VL.

GGT AGG AGG G-5' Ext : 5'-CCTCTGTCACA GTGCA CAA GAC ATC CAG ATG ACC CAG TCT CC \\
Bri : 3'-GG

AA-VL

-ApaLI-Q D

S

Ò

P S S +8 +9 +10

3. PCR

5'-GAC TGG GTG TAG TGA TCT AG-3' PCRpr.:

FR3) V * * S R D N S Y Y C A K

Bridge: 5'-G GTG TAG TCT AGA TCT CTG TTG AGA ... ATG ATA-5' A K

Ext : 3'-C CAC ATC ACT AGA TCT CTG TTG AGA ... ATG ATA-5' A K

-Xbal-

3'-XXX XXX XXX-VH

2. Ligation